## STIC Biotechnology Systems Branch

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- 1. EFS-Bio (<http://www.uspto.gov/ebc/efs/downloads/documents.htm>, EFS Submission User Manual ePAVE)
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   Alexandria, VA 22314

Revised 01/24/05



IFW16

RAW SEQUENCE LISTING DATE: 04/21/2005 PATENT APPLICATION: US/10/089,027 TIME: 11:37:43 Input Set : A:\NEB-166-PUS.APP.txt Output Set: N:\CRF4\04212005\J089027.raw 3 <110> APPLICANT: JACK, WILLIAM E. GARDNER, ANDREW BUZBY, PHILIP R. DiMEO, JAMES J. NEW ENGLAND BIOLABS, INC. 8 NEN LIFE SCIENCE PRODUCTS, INC. 10 <120> TITLE OF INVENTION: INCORPORATION OF MODIFIED NUCLEOTIDES BY ARCHAEON DNA POLYMERASES AND RELATED METHODS 13 <130> FILE REFERENCE: NEB-166-PUS 15 <140> CURRENT APPLICATION NUMBER: 10/089,027 16 <141> CURRENT FILING DATE: 2002-03-26 18 <150> PRIOR APPLICATION NUMBER: PCT/US00/26900 19 <151> PRIOR FILING DATE: 2000-09-29 21 <150> PRIOR APPLICATION NUMBER: 60/157,204 22 <151> PRIOR FILING DATE: 1999-09-30 Dees Not Comply 24 <160> NUMBER OF SEQ ID NOS: 33 26 <170> SOFTWARE: PatentIn Ver. 2.0 Corrected Diskette Needed ERRORED SEQUENCES 430 <210> SEQ ID NOT 31 431 <211> LENGTH: 631 432 <212> TYPE: DNA 433 <213> ORGANISM: Consensus using 9 degrees N or AmpliTaq 435 <220> FEATURE: 436 <221> NAME/KEY: misc feature 437 <222> LOCATION: (2)..(3) 438 <223> OTHER INFORMATION: n is a, c, g, or t 440 <220> FEATURE: 441 <221> NAME/KEY: misc\_feature 442 <222> LOCATION: (5)..(5) 443 <223> OTHER INFORMATION: s is g or c 445 <220> FEATURE: 446 <221> NAME/KEY: misc\_feature 447 <222> LOCATION: (11)..(12)

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453 <223> OTHER INFORMATION: n is a, c, g, or t

4

7

11

## RAW SEQUENCE LISTING DATE: 04/21/2005 PATENT APPLICATION: US/10/089,027 TIME: 11:37:43

Input Set : A:\NEB-166-PUS.APP.txt
Output Set: N:\CRF4\04212005\J089027.raw

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RAW SEQUENCE LISTING DATE: 04/21/2005 PATENT APPLICATION: US/10/089,027 TIME: 11:37:43

Input Set : A:\NEB-166-PUS.APP.txt
Output Set: N:\CRF4\04212005\J089027.raw

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     533 <223> OTHER INFORMATION: k is g or t
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     536 <221> NAME/KEY: misc feature
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     558 ccygggtacc gagctcngaa ttcngtaatc atggtcatag ctgtttcctt gtgtgaaatt
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     560 gttatccngc tcacaattcc acacaacata cngagccgga agcataaagt gtaaagcctg
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     562 gggtgcctaa tgagtgagct aactcacatt aattgcgttg cgctcacttg cccgctttcc
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     564 agtcgggaaa cctgtcgtgc cagctgcatt aatgaatcgg ccggagaggc ggtttgcgta
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     566 ttgggcgcca gggtggtttt tcttttcacc agtgagacgg gcaacagctg attgcccttc
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     568 accgcctggc cytgagagag ttgcagcaag cggtccacgc tggtttgccc cagcaggcga
                                                                                420
     570 aaatatggtg gttccgaaat cggcaaaatc ccttataaat caaaagaata gccccgagat
                                                                                480
     572 agggttgaag tgttgttcca gtttggaaca agagtccmct attaaagaam gtggactcca
                                                                                540
     574 acgkcaaagg gcgaaaaacc gtctwtcagg ggcgatggcc actacgkkaa ccwtcmccta
                                                                                600
E--> 576 atcaagtttt tggggkcgag kggccgttaa gccta
                                                                                 error
explanation
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## VERIFICATION SUMMARYDATE: 04/21/2005PATENT APPLICATION: US/10/089,027TIME: 11:37:44

Input Set : A:\NEB-166-PUS.APP.txt
Output Set: N:\CRF4\04212005\J089027.raw

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M:341 Repeated in SeqNo=31

L:576 M:252 E: No. of Seq. differs, <211> LENGTH:Input:631 Found:635 SEQ:31

L:626 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:32 after pos.:0

M:341 Repeated in SeqNo=32

L:861 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:33 after pos.:0

M:341 Repeated in SeqNo=33